



chain nodes :

7 8 9 10 11 12 16 22 23 24 25 26 27 28 29 30 31 32 33 35

ring nodes :

1 2 3 4 5 6 17 18 19 20 21

chain bonds :

1-21 2-7 4-8 5-32 6-31 8-9 8-16 10-12 10-11 17-22 17-29 18-23 18-30 19-24
19-25 21-35 24-27 25-26 26-33 27-28

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 17-18 17-21 18-19 19-20 20-21

exact/norm bonds :

1-2 1-6 1-21 2-3 2-7 3-4 4-5 4-8 5-6 8-16 10-12 10-11 17-18 17-21 17-22
18-19 18-23 19-20 19-24 20-21 24-27 26-33 27-28

exact bonds :

5-32 6-31 8-9 17-29 18-30 19-25 21-35 25-26

G1:0,S,N,C

G2:H,[*1]

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS
11:CLASS 12:CLASS 16:CLASS 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:CLASS
23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS
32:CLASS 33:CLASS 35:CLASS

=> d his

(FILE 'HOME' ENTERED AT 13:39:22 ON 21 SEP 2004)

FILE 'REGISTRY' ENTERED AT 13:39:30 ON 21 SEP 2004

L1 STRUCTURE UPLOADED

L2 9 S L1 SSS SAM

L3 104 S L1 SSS FULL

FILE 'CAPLUS, MEDLINE' ENTERED AT 13:41:34 ON 21 SEP 2004

L4 3 S L3

L5 3 DUP REM L4 (0 DUPLICATES REMOVED)